

Fig. 5B

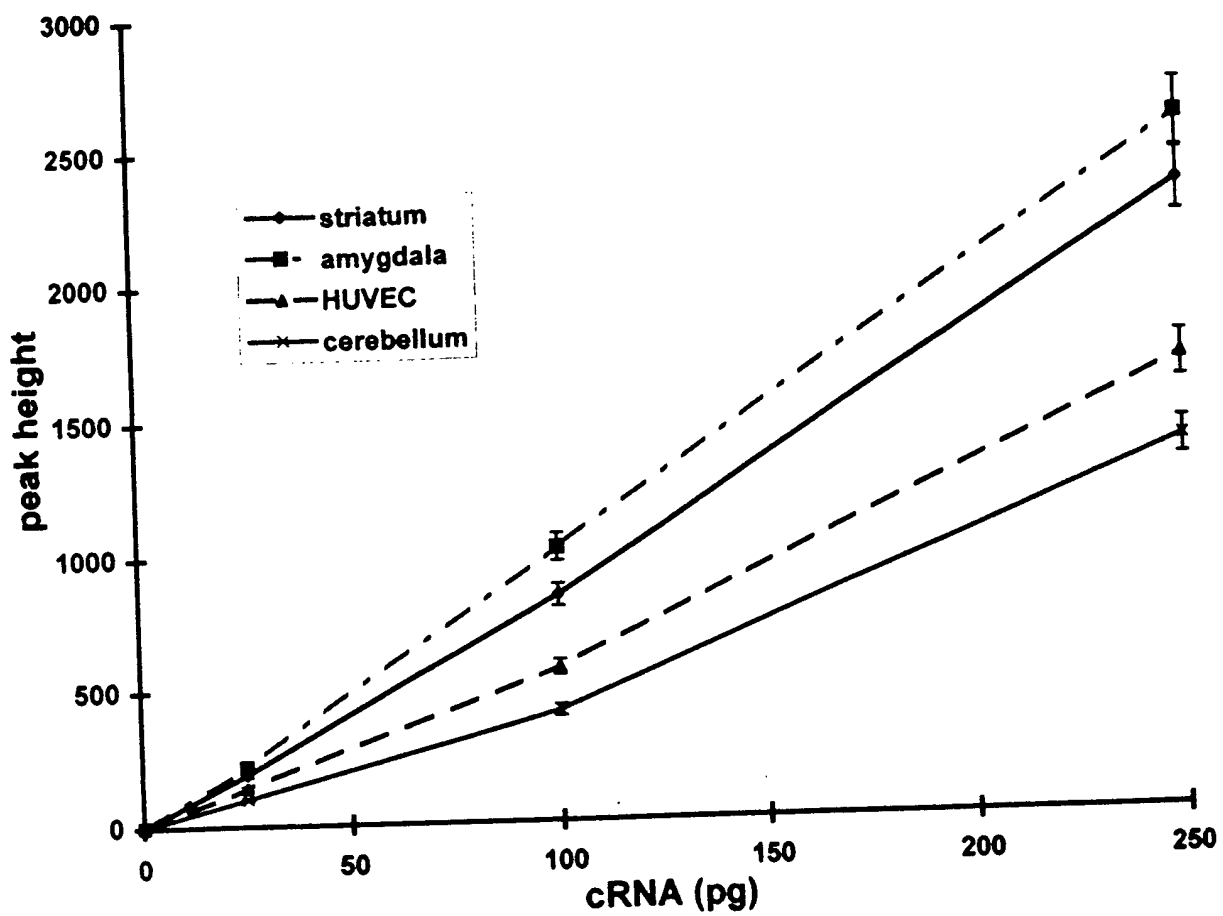


Fig. 6

[illegible]

B. pBS SK⁺/ DGT2 Multiple Cloning Site

C. pBS SK⁺/ DGT3 Multiple Cloning Site

Fig. 7

D. pBC SK⁺/ DGT4 Multiple Cloning Site

NotI BglII EcoRV BamH SacI BglII XhoI
5'-GCGGCCGCATAGATCT GATATC GGATCC TCACCACAGAGCTC AGTGAGAGAGATCTCTCGAG -3'
3'-CGCCGGCGTATCTAGA CTATAG CCTAGG AGTGGTGTCTCGAG TCAC TCTCTCTAGAGAGCTC-5'

E. pBS SK⁺/ DGT5 Multiple Cloning Site

NotI NcoI EcoRV SpHI Sall NcoI XhoI
5'-GCGGCCGCATCCATGG GATATC GCATGC TCACCACAGTCGAC AGTG AGAGCCATGG CTCGAG -3'
3'-CGCCGGCGTAGGTACC CTATAG CGTACG AGTGGTGTGAGCTG TCAC TCTC GGTACC GAGCTC-5'

Fig. 7 continued

0912669-10498

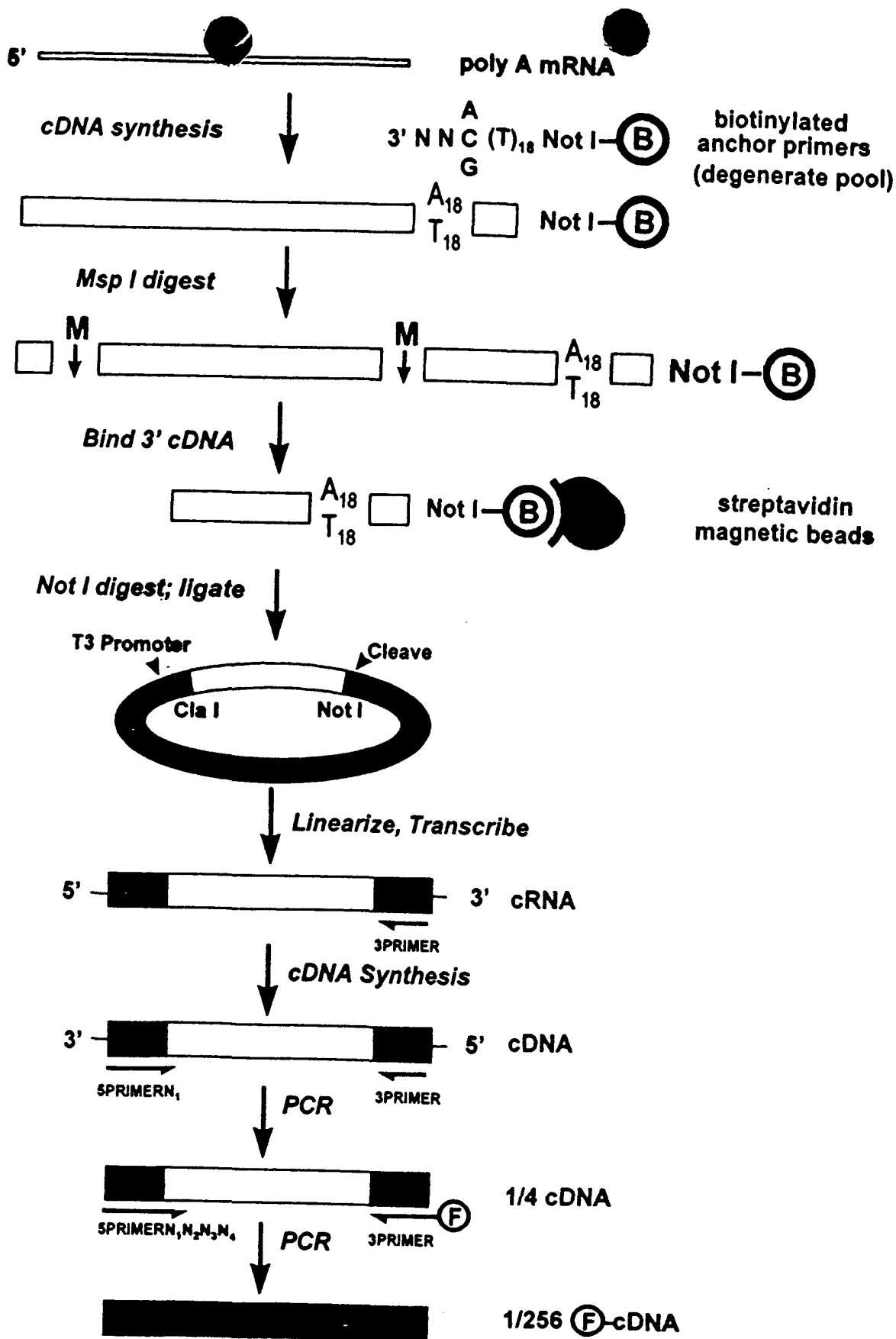


Fig. 8

09186869-110498